

FIGURE 1A

SEQ ID NO: 35	MW-----LLLTAAVISTLGTTHGLFGKIHPTNPEVTMNISQMITYWGYPAE	**
SEQ ID NO: 36	MWR-----FAVFLAAFFVQDVVSHG-----DPELHMTTPQIIERWGYPPAM	**
SEQ ID NO: 14	MAMAGHAPGGALPL-ILLVSCCGRIVSGASPAALRRVSGSGGLCDQLLLPLGYPCT	**
SEQ ID NO: 18	MALLGLMSFAALTFLVLT--VPRQAHASSRGNLGRNINPSVYGICASSVIVHGKQCQ	**
SEQ ID NO: 20	MRIQRFLATLATVTSILLGNPNVQCFDGGSHQKQH-----SLCEELIIPYGYPCS	**
	1	60
SEQ ID NO: 37	EYEVVTEDEGYIIGIDRIPYGRKNSE-NIGRR-PVAFIQHGLLASATNWNISNLPNNSLAFI	+
SEQ ID NO: 38	IYTVATDDGYIILEMHRIPFGKTNVTPNGKR-PVVFQHGGLLCASSDWVNLDPQSAGFL	+
SEQ ID NO: 14	EHNVEKDKGFLLSLQHIPHGKN--KAADSTGPPVFLQHGFLQGCGDTWFINSAEQSLGYI	+
SEQ ID NO: 18	EHEVTTDDGYIISLQRIPEGRG-KSSGSGTRKQPVVIOHGVLDGMTWLLNPPEQDLPLI	+
SEQ ID NO: 20	EHTIQTKDGFLLGLQVRVSSSSSLRLRNHGDGPPVILLHGLFMAGDAWFLNTPEQSLGFI	+
	61	120
SEQ ID NO: 37	LADAGYDVWLGNRSGNTWARRNLYYSPDSVEFWAFSDEMAYDLPATIDFILKKTGQDK	+
SEQ ID NO: 38	FADAGFDVWLGNMRGNTYSMKHKDLKPSHSAFWDWSWDEMATYDLNAMINHVLEVTGQDS	+
SEQ ID NO: 14	LADNGFDVWIGNVRGTRWSKGHSTFSVHDKLFDWDSWQELAEYDLLAMLGYYTVVTQS-K	+
SEQ ID NO: 18	LADNGFDVWIANTRGTRYRRHI SLDPSSQAYWNWSWDELVSDFPAVFNYVFSQTGQ-K	+
SEQ ID NO: 20	LADHGFDVWVGNVRGTRWSHGHI SLLEKKKQFWDWSWQELALYDVAEMINYINSVTNS-K	+
	121	180

[illegible]

FIGURE 1C

SEQ ID NO: 37	V--WNGGNDLLADPHDVDLLLSKL-PNLIYHRK---IPPNHLDFIWAMDAPQAVYNEIV	+	+	*	*	*	*	*	+	+	*
SEQ ID NO: 38	YLYWSDADWLADTPDVPDYLLTRINPAIVAQNN--HLPDYNHLDFWTGIRAPDDIYRPAI										
SEQ ID NO: 14	IWMGYGGDLADAVTDVQRTIREL---GSTPELLYIGDYGHIDFVMSVKAKDDVYVDLI										
SEQ ID NO: 18	LFISYGGRDALSDVRDVENLLDKLKFHDENKRSVQFIQEYAHADYIMGFNAKDLVYNVAVL										
SEQ ID NO: 20	IWMAYGGNDALADITDFQHTLKEI-----PSPPEVVYLENYGHVDFILSLQAKQDLYDPMI										
	361										420

SEQ ID NO: 37	SMMGTD-----NK	+
SEQ ID NO: 38	KLCTDD--YLGK	
SEQ ID NO: 14	RFLRENGWHNSY	
SEQ ID NO: 18	SFFNHQ-----V	
SEQ ID NO: 20	SFFKSSGKFSSM	
	421	433